IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Atty. Docket No: 03495-0213

In re patent application of

FLAMAND, MARIE *et al.*

Serial No. 09/980,839

Filed: December 7, 2001

For: EARLY DETECTION OF FLAVIVIRUSES USING THE NS1 GLYCOPROTEIN

STATEMENT TO SUPPORT FILING AND SUBMISSION IN ACCORDANCE WITH 37 C.F.R. §§ 1.821-1.825

Assistant Commissioner for Patents
Washington, D.C. 20231
Box SEQUENCE

Sir:

In connection with a Sequence Listing submitted concurrently herewith, the undersigned hereby states that:

1. the submission, filed herewith in accordance with 37 C.F.R. § 1.821(g), does not include new matter;

2. the content of the attached paper copy and the attached computer readable copy of the Sequence Listing, submitted in accordance with 37 C.F.R. § 1.821(c) and (e), respectively, are the same; and

3. all statements made herein of their own knowledge are true and that all statements made on information and belief are believed to be true; and further, that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United

States Code and that such willful false statements may jeopardize the validity of the application or any patent resulting therefrom.

Respectfully submitted,

April 23, 2002
Date

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SEQUENCE LISTING

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ALCON, SOPHIE
TALARMIN, ANTOINE
DESPRES, PHILIPPE
DEUBEL, VINCENT

<120> EARLY DETECTION OF FLAVIVIRUSES USING THE NS1 GLYCOPROTEIN

<130> 03495-0213

<140> 09/980,839
<141> 2001-12-07

<150> PCT/FR00/01620
<151> 2000-06-09

<150> FR 99/07290
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<150> FR 99/07361
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ata tca aat gaa ttg aac cac att cta ctt gaa aat gac atg aaa ttc   336
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gga aaa gcc aag atc ata gga gca gac aca cag aat acc acc ttc atc   480
Gly Lys Ala Lys Ile Ile Gly Ala Asp Thr Gln Asn Thr Thr Phe Ile
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atc gac ggc cca gac act cca gaa tgc ccc gat gac caa aga gcg tgg   528
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Asn Ile Trp Glu Val Glu Asp Tyr Gly Phe Gly Ile Phe Thr Thr Asn
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ata tgg ctg aaa ttg cgt gac tcc tac acc caa atg tgt gac cac cgg   624
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cta atg tca gct gcc gtc aag gac agc aag gca gtc cat gct gac atg   672
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ggg tac tgg ata gaa agt gaa aag aac gag acc tgg aag cta gcg aga   720
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tat gga gga cca ata tct cag cac aat tac aga cca ggg tat ttc aca   864
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305                 310                 315                 320

ggt cca tct ctc aga act aca aca gtc aca gga aag ata atc cat gaa  1008
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                325                 330                 335

tgg tgt tgc aga tcc tgc acg tta ccc ccc tta cgc ttc aga gga gaa  1056
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gac gga tgt tgg tat ggc atg gaa atc aga cca gtt aag gag aag gag  1104
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Thr Leu Tyr Leu Gly Val Met Val Gln Ala Asp Ser Gly Cys Val Ile
             20                  25                  30

Asn Trp Lys Gly Arg Glu Leu Lys Cys Gly Ser Gly Ile Phe Val Thr
         35                  40                  45

Asn Glu Val His Thr Trp Thr Glu Gln Tyr Lys Phe Gln Ala Asp Ser
     50                  55                  60

Pro Lys Arg Leu Ser Ala Ala Ile Gly Lys Ala Trp Glu Glu Gly Val
 65                  70                  75                  80

Cys Gly Ile Arg Ser Ala Thr Arg Leu Glu Asn Ile Met Trp Lys Gln
                 85                  90                  95

Ile Ser Asn Glu Leu Asn His Ile Leu Leu Glu Asn Asp Met Lys Phe
            100                 105                 110

Thr Val Val Val Gly Asp Ala Asn Gly Ile Leu Ala Gln Gly Lys Lys
        115                 120                 125

Met Ile Arg Pro Gln Pro Met Glu His Lys Tyr Ser Trp Lys Ser Trp
    130                 135                 140
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Gly Lys Ala Lys Ile Ile Gly Ala Asp Thr Gln Asn Thr Thr Phe Ile
145                 150                 155                 160

Ile Asp Gly Pro Asp Thr Pro Glu Cys Pro Asp Asp Gln Arg Ala Trp
                165                 170                 175

Asn Ile Trp Glu Val Glu Asp Tyr Gly Phe Gly Ile Phe Thr Thr Asn
            180                 185                 190

Ile Trp Leu Lys Leu Arg Asp Ser Tyr Thr Gln Met Cys Asp His Arg
        195                 200                 205

Leu Met Ser Ala Ala Val Lys Asp Ser Lys Ala Val His Ala Asp Met
    210                 215                 220

Gly Tyr Trp Ile Glu Ser Glu Lys Asn Glu Thr Trp Lys Leu Ala Arg
225                 230                 235                 240

Ala Ser Phe Ile Glu Val Lys Thr Cys Ile Trp Pro Lys Ser His Thr
                245                 250                 255

Leu Trp Ser Asn Gly Val Leu Glu Ser Glu Met Ile Ile Pro Lys Ile
            260                 265                 270

Tyr Gly Gly Pro Ile Ser Gln His Asn Tyr Arg Pro Gly Tyr Phe Thr
        275                 280                 285

Gln Thr Ala Gly Pro Trp His Leu Gly Lys Leu Glu Leu Asp Phe Asp
    290                 295                 300

Leu Cys Glu Gly Thr Thr Val Val Val Asp Glu His Cys Gly Asn Arg
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Gly Pro Ser Leu Arg Thr Thr Thr Val Thr Gly Lys Ile Ile His Glu
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Trp Cys Cys Arg Ser Cys Thr Leu Pro Pro Leu Arg Phe Arg Gly Glu
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Asp Gly Cys Trp Tyr Gly Met Glu Ile Arg Pro Val Lys Glu Lys Glu
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